



# SEQUENCE LISTING

<110> Sera, Takashi

<120> Nuclear-Envelope and Nuclear-Lamina Binding Chimeras for Modulating Gene Expression

<130> 109845-163

<160> 21

<170> PatentIn version 3.3

<210> 1

<211> 25

<212> PRT

<213> Artificial

<220>

<223> Zinc finger domain

<220>

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<222> (2)..(5)

<223> Amino acids 2-5 are Xaa wherein Xaa = any amino acid, and up to two amino acids can be missing.

<220>

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<222> (7)..(18)

<223> Xaa can be any amino acid

<220>

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<222> (20)..(24)

<223> Amino acids 20-24 are Xaa wherein Xaa = any amino acid, and up to two amino acids can be missing.

<400> 1

Cys Xaa Xaa Xaa Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa

1 5 10 15

Xaa Xaa His Xaa Xaa Xaa Xaa Xaa His

20 25

<210> 2

<211> 32

<212> PRT

<213> Artificial

<220>

<223> Second zinc finger domain

<220>

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<222> (1)..(3)

<223> Xaa can be any amino acid

<220>

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<222> (5)..(8)

<223> Amino acids 5-8 are Xaa wherein Xaa = any amino acid, and up to two amino acids can be missing

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<220>
<221> MISC_FEATURE
<222> (10)..(14)
<223> Xaa can be any amino acid

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<222> (15)..(15)
<223> Amino acid 15 is Z(-1) wherein Z(-1) = Arg, Lys, Gln, Asn, Thr,
Met, Leu, Ile, Glu or Asp.

<220>
<221> MISC_FEATURE
<222> (16)..(16)
<223> Xaa can be any amino acid

<220>
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<222> (17)..(17)
<223> Amino acid 17 is Z2 wherein Z2 = Ser, Arg, Asn, Gln, Thr, Val,
Ala, Asp or Glu.

<220>
<221> MISC_FEATURE
<222> (18)..(18)
<223> Amino acid 18 is Z3 wherein Z3 = His, Lys, Asn, Gln, Ser, Ala,
Val, Thr, Asp, or Glu

<220>
<221> MISC_FEATURE
<222> (19)..(20)
<223> Xaa can be any amino acid

<220>
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<222> (21)..(21)
<223> Amino acid 21 is Z6 wherein Z6 = Arg, Lys, Gln, Asn, Thr, Tyr,
Leu, Ile, Met, Glu or Asp.

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<221> MISC_FEATURE
<222> (23)..(27)
<223> Amino acids 23-27 are Xaa wherein Xaa = any amino acid, adn up to
two amino acids can be missing.

<220>
<221> MISC_FEATURE
<222> (29)..(32)
<223> Xaa can be any amino acid

<400> 2

Xaa Xaa Xaa Cys Xaa Xaa Xaa Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa
1 5 10 15

Xaa Xaa Xaa Xaa Xaa His Xaa Xaa Xaa Xaa Xaa His Xaa Xaa Xaa Xaa
20 25 30

<210> 3
<211> 28
<212> PRT
<213> Artificial

<220>

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<223> Zinc finger domain

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<222> (13)..(13)

<223> Amino acid 13 is Z(-1) wherein Z(-1) = Arg, Lys, Gln, Asn, Thr, Met, Leu, Ile, Glu or Asp.

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<221> MISC\_FEATURE

<222> (15)..(15)

<223> Amino acid 15 is Z2 wherein Z2 = Ser, Arg, Asn, Gln, Thr, Val, Ala, Asp or Glu.

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<221> MISC\_FEATURE

<222> (16)..(16)

<223> Amino acid 16 is Z3 wherein Z3 = His, Lys, Asn, Gln, Ser, Ala, Val, Thr, Asp or Glu.

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<221> MISC\_FEATURE

<222> (19)..(19)

<223> Amino acid 19 is Z6 wherein Z6 = Arg, Lys, Gln, Asn, Thr, Tyr, Leu, Ile, Met, Glu or Asp.

<400> 3

Pro Tyr Lys Cys Pro Glu Cys Gly Lys Ser Phe Ser Xaa Ser Xaa Xaa  
1 5 10 15

Leu Gln Xaa His Gln Arg Thr His Thr Gly Glu Lys  
20 25

<210> 4

<211> 5

<212> PRT

<213> Artificial

<220>

<223> Synthetic flexible linker peptide for linking together multi-finger zinc finger domains

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Gly Gly Gly Gly Ser  
1 5

<210> 5

<211> 11

<212> PRT

<213> Human immunodeficiency virus

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<221> MISC\_FEATURE

<222> (1)..(11)

<223> HIV Tat protein domain

<400> 5

Tyr Gly Arg Lys Lys Arg Arg Gln Arg Arg Arg  
1 5 10

<210> 6  
<211> 9  
<212> DNA  
<213> Human immunodeficiency virus

<220>  
<221> misc\_feature  
<222> (1)..(9)  
<223> HIV DNA Binding Domain

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gcagaagcc

9

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<211> 19  
<212> DNA  
<213> Artificial

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<223> DNA target sequence

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gtgtgggtga gtgagtgtg

19

<210> 8  
<211> 19  
<212> DNA  
<213> Artificial

<220>  
<223> DNA target sequence

<400> 8  
ggggctgggg gcggtgtct

19

<210> 9  
<211> 7  
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<213> Simian virus 40

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<222> (1)..(7)  
<223> Peptide from SV40 large T antigen

<400> 9

Pro Lys Lys Lys Arg Lys Val  
1 5

<210> 10  
<211> 16  
<212> PRT  
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<220>  
<223> Peptide, residues 43-58 of the Antennapeida homeodomain protein

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<221> MISC\_FEATURE

<222> (1)..(16)  
<223> Peptide, residues 43-58 of the Antennapeida homeodomain protein

<400> 10

Arg Gln Ile Lys Ile Trp Phe Gln Asn Arg Arg Met Lys Trp Lys Lys  
1 5 10 15

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<213> Herpes Simplex Virus

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<222> (1)..(34)  
<223> Residues 267-300 of the HSV VP22 protein

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Asp Ala Ala Thr Ala Thr Arg Gly Arg Ser Ala Ala Ser Arg Pro Thr  
1 5 10 15

Glu Arg Pro Arg Ala Pro Ala Arg Ser Ala Ser Arg Pro Arg Arg Pro  
20 25 30

Val Glu

<210> 12  
<211> 11  
<212> PRT  
<213> Artificial

<220>  
<223> Synthetic peptide modeled after the protein transduction domain  
of the human immunodeficiency virus TAT protein having cellular  
uptake activity

<400> 12

Tyr Ala Arg Ala Ala Ala Arg Gln Ala Arg Ala  
1 5 10

<210> 13  
<211> 9  
<212> PRT  
<213> Artificial

<220>  
<223> Synthetic peptide modeled after the protein transduction domain  
of the human immunodeficiency virus TAT protein having cellular  
uptake activity, referred to as "R9"

<400> 13

Arg Arg Arg Arg Arg Arg Arg Arg  
1 5

<210> 14  
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<212> PRT  
<213> Artificial

<220>  
<223> D-penetratin peptide

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<222> (1)..(16)  
<223> All amino acids are in the D-form

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Arg Gln Ile Lys Ile Trp Phe Gln Asn Arg Arg Met Lys Trp Lys Lys  
1 5 10 15

<210> 15  
<211> 16  
<212> PRT  
<213> Artificial

<220>  
<223> Peptide Syn B1 from Antennapedia homeodomain protein

<400> 15

Arg Gly Gly Arg Leu Ser Tyr Ser Arg Arg Arg Phe Ser Thr Ser Thr  
1 5 10 15

<210> 16  
<211> 10  
<212> PRT  
<213> Artificial

<220>  
<223> L-SynB3 peptide from Antennapedia homeodomain protein

<400> 16

Arg Arg Leu Ser Tyr Ser Arg Arg Arg Phe  
1 5 10

<210> 17  
<211> 10  
<212> PRT  
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<220>  
<223> D-SynB3 peptide from Antennapedia homeodomain pro

<220>  
<221> MISC\_FEATURE  
<222> (1)..(10)  
<223> All amino acids are in the D-form

<400> 17

Arg Arg Leu Ser Tyr Ser Arg Arg Arg Phe  
1 5 10

<210> 18  
<211> 8

<212> PRT  
<213> Artificial

<220>  
<223> Flag Epitope Peptide

<400> 18

Asp Tyr Lys Asp Asp Asp Asp Lys  
1 5

<210> 19  
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<212> PRT  
<213> Artificial

<220>  
<223> Artificial peptide linker

<400> 19

Gly Gly Gly Gly Ser  
1 5

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<212> PRT  
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Gly Gly Gly Ser  
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<220>  
<223> Artificial peptide linker

<400> 21

Gly Gly Ser  
1